

SEQUENCE LISTING

<110> Pang, Danny Z.
<120> Use of decorin in a cosmetic or dermatological composition
<160> 6
<170> PatentIn ver. 2.0

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TRANSLATE = PROTEIN

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gag ccc tcc cta ggc cca gtg tgc ccc ttc cgc tgt caa tgc cat ctt 96
Glu Pro Ser Leu Gly Pro Val Cys Pro Phe Arg Cys Gln Cys His Leu
20 25 30

cga gtg gtc cag tgt tct gat ttg ggt ctg gac aaa gtg cca aag gat 144
Arg Val Val Gln Cys Ser Asp Leu Gly Leu Asp Lys Val Pro Lys Asp
35 40 45

ctt ccc cct gac aca act ctg cta gac ctg caa aac aac aaa ata acc 192
Leu Pro Pro Asp Thr Thr Leu Leu Asp Leu Gln Asn Asn Lys Ile Thr
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gaa atc aaa gat gga gac ttt aag aac ctg aag aac ctt cac gca ttg 240
Glu Ile Lys Asp Gly Asp Phe Lys Asn Leu Lys Asn Leu His Ala Leu
65 70 75 80

att ctt gtc aac aat aaa att agc aaa gtt agt cct gga gca ttt aca 288
Ile Leu Val Asn Asn Lys Ile Ser Lys Val Ser Pro Gly Ala Phe Thr
85 90 95

cct ttg gtg aag ttg gaa cga ctt tat ctg tcc aag aat cag ctg aag 336
Pro Leu Val Lys Leu Glu Arg Leu Tyr Leu Ser Lys Asn Gln Leu Lys
100 105 110

gaa ttg cca gaa aaa atg ccc aaa act ctt cag gag ctg cgt gcc cat		384	
Glu Leu Pro Blu Lys Met Pro Lys Thr Leu Gln Glu Leu Arg Ala His			
115	120	125	
gag aat gag atc acc aaa gtg cga aaa gtt act ttc aat gga ctg aac		432	
Glu Asn Glu Ile Thr Lys Val Arg Lys Val Thr Phe Asn Gly leu Asn			
130	135	140	
cag atg att gtc ata gaa ctg ggc acc aat ccg ctg aag agc tca gga		480	
Gln Met Ile Val Ile Glu Leu Gly Thr Asn Pro Leu Lys Ser Ser Gly			
145	150	155	160
att gaa aat ggg gct ttc cag gga atg aag aag ctc tcc tac atc cgc		528	
Ile Glu Asn Gly Ala Phe Gln Gly Met Lys Lys Leu Ser Tyr Ile Arg			
165	170	175	
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Ile Ala Asp Thr Asn Ile Thr Ser Ile Pro Gln Gly Leu Pro Pro Ser			
180	185	190	
ctt acg gaa tta cat ctt gat ggc aac aaa atc agc aga gtt gat gca		624	
Leu Thr Glu Leu His Leu Asp Gly Asn Lys Ile Ser Arg Val Asp Ala			
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gct agc ctg aaa gga ctg aat aat ttg gct aag ttg gga ttg agt ttc		672	
Ala Ser Leu Lys Gly Leu Asn Asn Leu Ala Lys Leu Gly Leu Ser Phe			
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Asn Ser Ile Ser Ala Val Asp Asn Gly Ser Leu Ala Asn Thr Pro His			
225	230	235	240
ctg agg gag ctt cac ttg gac aac aac aag ctt acc aga gta cct ggt		768	
Leu Arg Glu Leu His Leu Asp Asn Asn Lys Leu Thr Arg Val Pro Gly			
245	250	255	
ggg ctg gca gag cat aag tac atc cag gtt gtc tac ctt cat aac aac		816	
Gly Leu Ala Glu His Lys Tyr Ile Gln Val Val Tyr Leu His Asn Asn			
260	265	270	
aat atc tct gta gtt gga tca agt gac ttc tgc cca cct gga cac aac		864	
Asn Ile Ser Val Val Gly Ser Ser Asp Phe Cys Pro Pro Gly His Asn			
275	280	285	
acc aaa aag gct tct tat tcg ggt gtg agt ctt ttc agc aac ccg gtc		912	
Thr Lys Lys Ala Ser Tyr Ser Gly Val Ser Leu Phe Ser Asn Pro Val			
290	295	300	
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Gln Tyr Trp Glu Ile Gln Pro Ser Thr Phe Arg Cys Val Tyr Val Arg			
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35 40 45

Leu Pro Pro Asp Thr Thr Leu Leu Asp Leu Gln Asn Asn Lys Ile Thr
50 55 60

Glu Ile Lys Asp Gly Asp Phe Lys Asn Leu Lys Asn Leu His Ala Leu
65 70 75 80

Ile Leu Val Asn Asn Lys Ile Ser Lys Val Ser Pro Gly Ala Phe Thr
85 90 95

Pro Leu Val Lys Leu Glu Arg Leu Tyr Leu Ser Lys Asn Gln Leu Lys
100 105 110

Glu Leu Pro Glu Lys Met Pro Lys Thr Leu Gln Glu Leu Arg Ala His
115 120 125

Glu Asn Glu Ile Thr Lys Val Arg Lys Val Thr Phe Asn Gly Leu Asn
130 135 140

Gln Met Ile Val Ile Glu Leu Gly Thr Asn Pro Leu Lys Ser Ser Gly
145 150 155 160

Ile Glu Asn Gly Ala Phe Gln Gly Met Lys Lys Leu Ser Tyr Ile Arg
165 170 175

Ile Ala Asp Thr Asn Ile Thr Ser Ile Pro Gln Gly Leu Pro Pro Ser
180 185 190

Leu Thr Glu Leu His Leu Asp Gly Asn Lys Ile Ser Arg Val Asp Ala
195 200 205

Ala Ser Leu Lys Gly Leu Asn Asn Leu Ala Lys Leu Gly Leu Ser Phe
210 215 220

Asn Ser Ile Ser Ala Val Asp Asn Gly Ser Leu Ala Asn Thr Pro His
225 230 235 240

Leu Arg Glu Leu His Leu Asp Asn Asn Lys Leu Thr Arg Val Pro Gly
245 250 255

Gly Leu Ala Glu His Lys Tyr Ile Gln Val Val Tyr Leu His Asn Asn
260 265 270

Asn Ile Ser Val Val Gly Ser Ser Asp Phe Cys Pro Pro Gly His Asn
275 280 285

Thr Lys Lys Ala Ser Tyr Ser Gly Val Ser Leu Phe Ser Asn Pro Val
290 295 300

Gln Tyr Trp Glu Ile Gln Pro Ser Thr Phe Arg Cys Val Tyr Val Arg
305 310 315 320

Ser Ala Ile Gln Leu Gly Asn Tyr Lys
325